Ref #	Hit	s Searc	ch Query	DBs	Default Operator	Plurals	Time Stamp
L1		7   "100	2862".pn.	EPO; DERWENT	OR	OFF	2005/02/01 12:30
L2	:	2 KIAA	near5 prostate	US-PGPUB; USPAT; EPO; DERWENT	OR	OFF	2005/02/01 12:31
L3	· · · · (	selad	lin-1 near5 prostate	US-PGPUB;	OR	OFF	2005/02/01 12:32
	r in			USPAT; EPO; DERWENT			
L4		3 selad	lin-1 and prostate	US-PGPUB; USPAT; EPO; DERWENT	OR	OFF	2005/02/01 12:33
<b>L5</b>		5 selad	lin-1	US-PGPUB; USPAT; EPO; DERWENT	OR	OFF	2005/02/01 12:33

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2 (bases 1 to 4882)
Ohara,O., Nagase,T., Kikuno,R. and Nomura,N.
Direct Submission
Submitted (24-NOV-1994) Osamu Ohara, Kazusa DN
1532-3, Yana, Kisarazu, Chiba 292-0812, Japan
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                                                                                                                    Nagase, T., Miyajima, N., Tanaka, A., Sazuka, T., Seki, N., Sato, S., Tabata, S., Ishikawa, K.-i., Kawarabayasi, Y., Kotani, H. and Nomura, N. Prediction of the coding sequences of unidentified human genes. III. The coding sequences of 40 new genes (KIAA00120) deduced by analysis of cDNA clones from human cell line KG-1 NNA Reg. 2 (1), 37,43.(1995)
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D43636
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On May 9, 2002 this sequence version replaced gi:598957
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ARAADSVLNGHRSKGLCDSAKKDDLPELAGPALSTVPPASLKOTASGRCCLFRUEEDE
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RRDSSEGPPGSEGDGGGGSKCSNASGGLVESCHASPSLHGTASGGSBSSGGGNPTNTSGTT
RRCAGPSNSMQLASRSAGELVESLKLMSLCLGSQLHGSTKYIIDPQNGLSFSSVKVQE
KSTWANGISSTGNAGQVPAVGGIKFFSDHMADTTTELERIKSKNLKNNVLQLPLCEKT
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derived from ha01240)."
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/note="KIAA0096
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Direct Submission
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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derived from a RT-PCR product (Lung) and 1014 - 4882 was
derived from ha01240)."
/gene="KIAA0096"
/note="KIAA0096 gene product is related to
kinase."
                                                   <1. .2302
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                                                                                                                                                                                                                                  _line="KG-1"
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Strausberg, R.L., Feingold, E.A., Grouse, L.H., Derge, J.G., Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D. Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K., Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Haieh, F., Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L., Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L., Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S., Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S., Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J.,
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                                                                                                                                                                                                                                                                                                                                                                  5140 bp
Homo sapiens SNF-1 related kinase,
IMAGE:4375623), complete cds.
BC071567
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Eukaryota;
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                                                                      Al-Mahmood,S., Colin,S. and Schneider,C. Genes involved in regulating angiogenesis, pharmaceutical preparations containing same and applications thereof Patent; WO 03074073-A 53 12-SEP-2003;
                                                                                                                                 Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                       Sequence 53
AX828407
                                                             Gene
                                                                                                                                                                   Homo sapiens (human)
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/db_xref="taxon:9606"
/noTe="EMBL/GenBank Accession No.
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/organism="Homo sapiens"
/mol_type="unassigned DN
/db_xref="taxon:9606"
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                                                                                                                                                                                              GI:39838407
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                                         Submitted (11-NOV-1992)
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nomura,N., Miyajima,N., Sazuka,T., Tanaka,A., Kawarabayasi,Y., Sato,S., Nagase,T., Seki,N., Ishikawa,K. and Tabata,S. Prediction of the coding sequences of unidentified human genes. I The coding sequences of 40 new genes (KIAA0001-KIAA0040) deduced
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            analysis of randomly sampled cDNA clones from human immature
                                                                                                                                3 (bases 1 to 4187)
Ohara,O., Nagase,T., Kikuno,R. and Nomura,N.
Direct Submission
                  1532-3, Yana, Kisarazu, Chiba 292-0812, Japan (B-mail:cdnainfo@kazusa.or.jp, Tel:+81-438-52-3913) (D. Dec 22, 1999 this sequence version replaced gi:285996. Sequence updated (20-Dec-1999)
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VfvClfflblslifdlyyyrawvyfklssaprlheggrvrliqqxw
CTGREGWTD15LRVGKKXTHKQIMIXIMMDILEVDTKKQIVRVEGVSKTFM
CTGRFGWLTVSLRVGKKXTHKQIMIXIMMDILEVDTKKQIVRVEFLVTMGQVTALTIS
IGWTLFVLFELDDLTVGGLIMGTG1ESSSHKYGLFQHICTAYELVLADGSFVRCTPSE
IGWTLFVLFELDDLTVGGLIMGTG1ESSSHKYGLFQHICTAYELVLADGSFVRCTPSE
IGWTLFVLFWGSCGTLGFLVAAEIRIIFAKKYVYLRFEFVRGLEAICAKETHESQRQEN
HFVEGLLYSLDEAVIMTGVMTDEAFESKLNSIGNYYKFWFFKHVENYLKTNRESGREEYI
PLRHYYHRHTRSIFWELQDIIFGNNDIFTRYLFGWMVFFKISLLKLTQGGTLRKLYED
HTVVDMLVFMKCLQOALHTFONDIHYFIMLGFFIFSQFGLVHFKGNEAELYIDIG
HTVQDMLVFMKCLQOALHTFONDIHYFIMLGFFIFSQFGLVHFKGNEAELYIDIG
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/codon_start=3
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/protein_id="BAA02806.2"
/db_xref="GI:6630632"
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/cell_type="myoblast"
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/mol_type="mRNA"
/db_xref="taxon:9606".
/clone="ha005171"
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ROGER NITSCH, ISABELL GREEVE
OS Homo sapiens (human)
PN JP 2002530076-A/1
PD 17-SEP-2002
PF 12-NOV-1999 JP 200058255:
PR 12-NOV-1998 EP 9812147/
PI ROGER NITSCH, ISABELL GREE
PC C12N15/09, A6IX38/00, A6IX6
PC A6IP1/00, A6IP15/00, A6IP5
PC A6IP25/28,
PC A6IP25/28,
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PC G01N33/53,G01N33/53,C12N15/00,C12N5/00,A61K37/0
of diagnosing or treating neurological diseases and
cel1
CC degeneration
FH Key Location/Qualifiers
FT Source 1. 4248
FT fource 1. 4248
FT fource 1. 4248
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                       AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA
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                                                                                                              AGAAGACACAAAAAACATCATGATCAACCTGATGGACAT 400
                                                                                                                                                                                       CCTTCATGTGCACGGGGGCCCCTGGCTGGCTCACTGTCTCACTACGTGTCGGGAAGTACA
                                                                                                                                                                                                                                 AGCAGCGCGTGCGGGACATCCAGAAGCAGGTGCGGGAATGGAAGGAGCAGGGTAGCAAGA
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12-NOV-1998 EP 98121478.6
ROGER NITSCH, ISABELL GREEVE
C12N15/09, A61K38/00, A61K39/395, A61K39/395, A61K45/00, A61K48/00,
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A61P11/00,A61P15/00,A61P21/00,A61P25/00,A61P25/14,A61P25/16,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 1.7e-67;
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Length 4248;

(human)'.

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Gaps

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DNA Res. 1 (1), 47-56 (1994)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUMRSC390 4187 bp mRNA linear Homo sapiens mRNA for KIAA0018 protein, partial cds.
                                                                                                                                                                                                                      Submitted (11-NOV-1992) Osamu Ohara, Kazusa DNA Research Institute; 1532-3, Yana, Kisarazu, Chiba 292-0812, Japan (E-mail:cdnainfo@kazusa.or.jp, Tel:+81-438-52-3913) On Dec 22, 1999 this sequence version replaced gi:285996. Sequence updated (20-Dec-1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           myeloid cell line KG-1
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Mammalia; Eutheria;.Primates; Catarrhini; Hominidae; Homo.
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/gene="KIAA0018"
<1. .1589
                                                    /cell_
                                                                                                          /db_xref="taxon:9606"
/clone="ha005171"
                                                                                                                                              organism="Homo sapiens"
/mol_type="mRNA"
                                                                                          'sex="male"
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                                                                                                                                                                                                    ocation/Qualifiers
                              ell_type="myoblast"
.4187
                                                                           line="KG-1"
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Strausberg, R.L., Feingold, E.A., Grouse, L.H., Derge, J.G., Klausmer, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D., Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K., Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,
                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 4202)
                                                                                                                                                                                                                                                                          Homo sapiens 24-dehydrocholesterol reductase, MGC:10569 IMAGE:3638639), complete cds.
                                                                                                                                                                                  Homo sapiens
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VFVCLFILPISIIFDIYYYVBAWVVFKLSSAPRLHBQRVRDIJQKOVBEMKEQGSKTFM
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NSDLFYAVPMSCGTLGFLVAAEIRIIPAKKYVKLRFEPVRGLEAICAKFTHESQRQEN
HFVEGLLYSLDEAVINTGVMTDEAEPSKLNSIGNYYKFWFFKHVENYLKTNREGLEYI
PLRHYYHRHTRSIFWELQDIIPFGNNPIFRYPIWLCFFILPSQPGLVHFKGNEAELYIDIG
PHVVQDMLVPWKCLQQALHTFQNDIHYYFIWLCFFILPSQPGLVHFKGNEAELYIDIG
AYGEFRVKHFEARSCMRQLEKFVRSVHGFQMLYADCYMNREBFWEMFDGSLYHKLREK
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/product="KIAA0018 protein"
/protein_id="BAA02806.2"
/db_xref="GI:6630632"
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/note="Start codon is not identified. ha00517 cDNA clone for KIAA0018 has a 1-bp deletion at the position 1167 of the sequence of KIAA0018."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LGCQDAFPEVYDKICKAARH"
1590. .>4187
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                                                                                                                                                                                                                                                   GI:13325123
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Pred. No. 9.2e-116;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      4187
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                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                    mRNA (cDNA clone
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